

pH-sensitive Gold Nanorods for Non-Steroidal Anti-Inflammatory Drugs (NSAIDs) Delivery and DNA-binding studies

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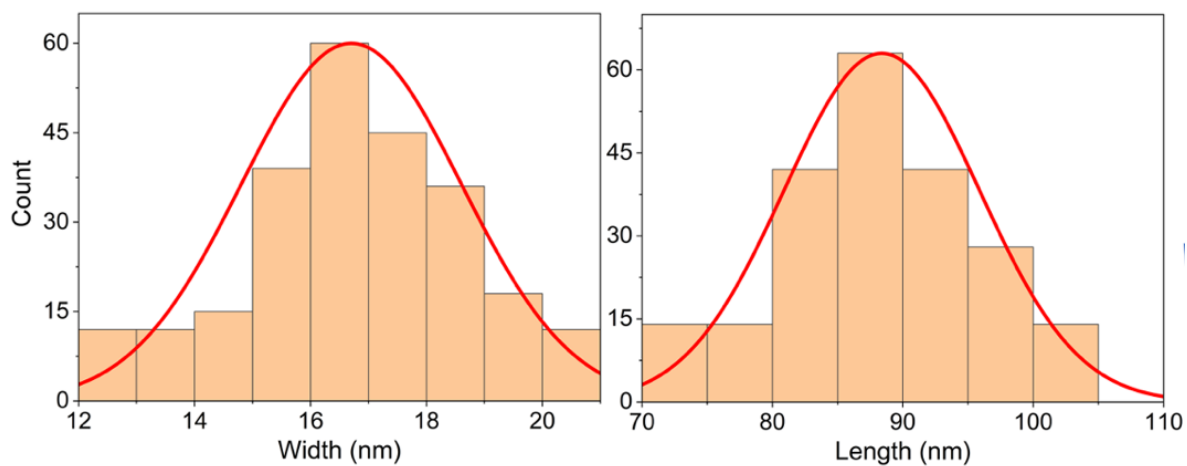


Figure S1. Gaussian distributions of sizes of AuNRs.

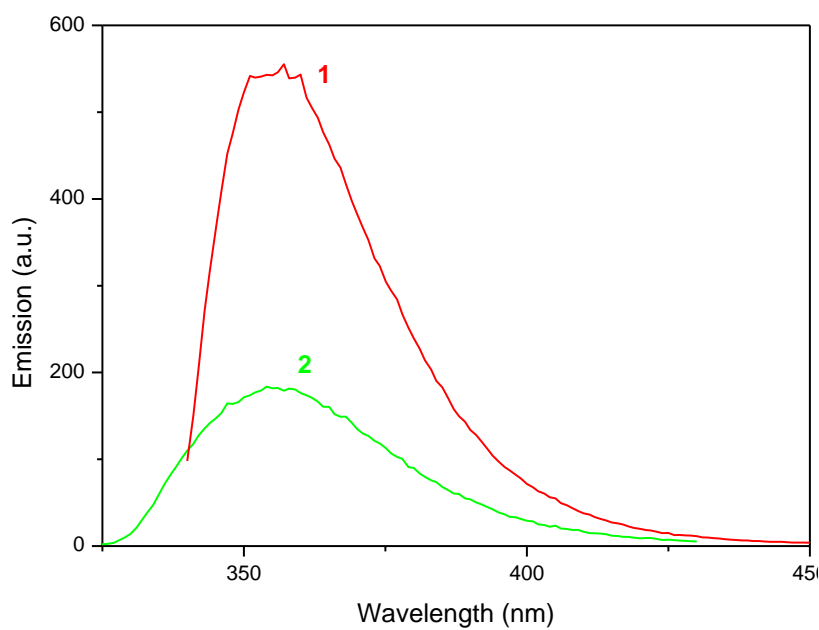


Figure S2. Comparison between the emission spectra of the free NAP solution (red line, $\lambda_{\text{max,em}} = 357 \text{ nm}$) and AuNRs@PEG@NAP ($\lambda_{\text{max,em}} = 359 \text{ nm}$).

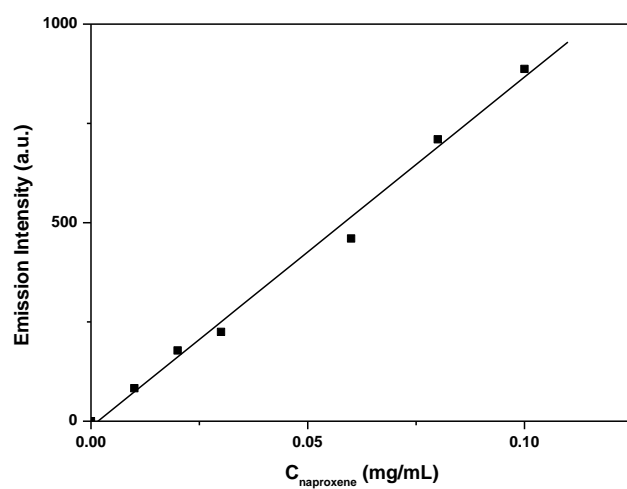
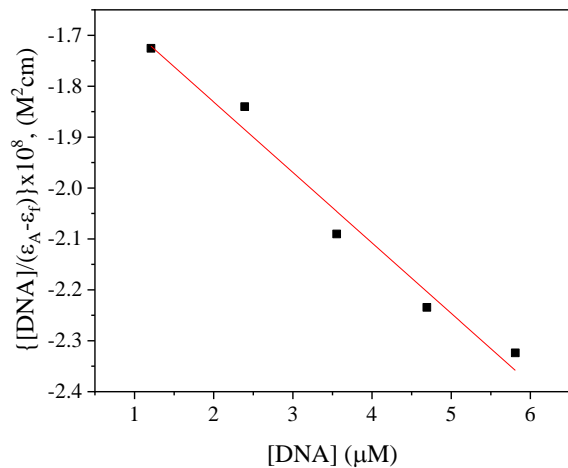


Figure S3. Calibration line of the emission intensity of NAP at 357 nm as a function of the concentration of NAP.

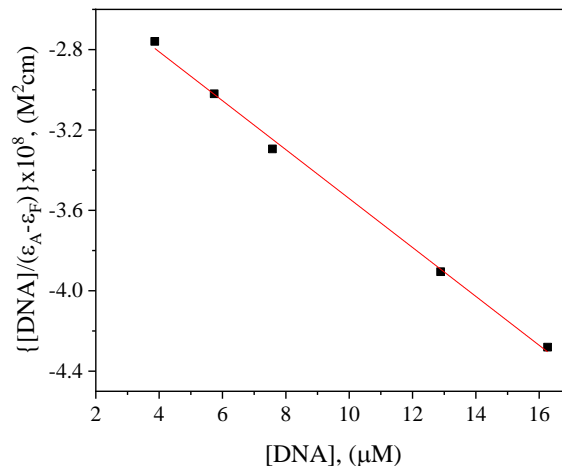
Linear Regression for Data1_B:

$$Y = A + B * X$$

Parameter	Value	Error	
A	-14.24415	18.75409	
B	8809.0301	339.18614	
<hr/>			
R	SD	N	P
0.99631	31.35015	7	<0.0001



AuNRs@PEG



AuNRs@PEG@NAP

Figure S4. Plot of $\frac{[DNA]}{(\epsilon_A - \epsilon_f)}$ versus [DNA] for AuNRs@PEG and AuNRs@PEG@NAP.

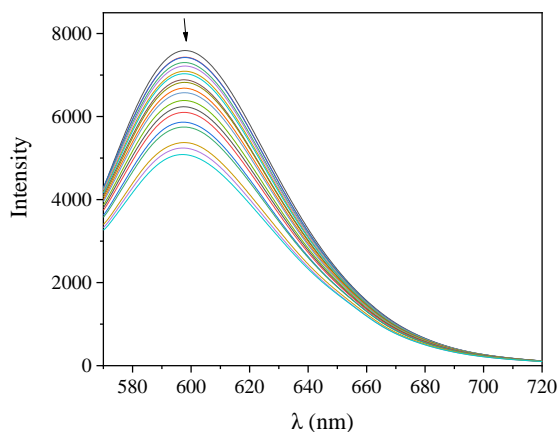
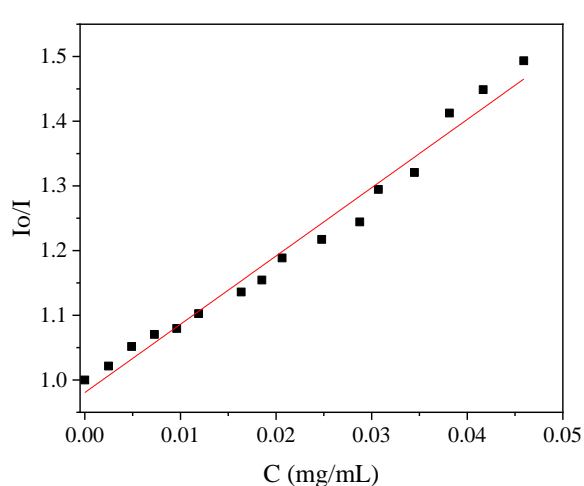
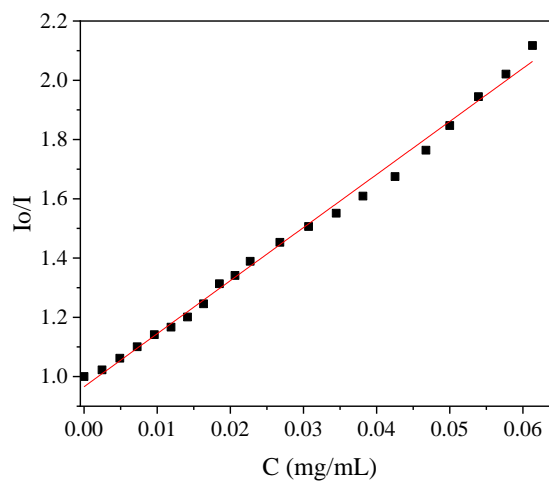


Figure S5. Fluorescence emission spectra ($\lambda_{\text{excitation}} = 540$ nm) for EB–DNA adduct ([EB] = 20 μM , [DNA] = 26 μM) in buffer solution (150 mM NaCl and 15 mM trisodium citrate at pH = 7.0) in the absence and presence of increasing amounts of AuNRs@PEG. The arrow shows the changes of intensity upon increasing amounts of AuNRs@PEG.

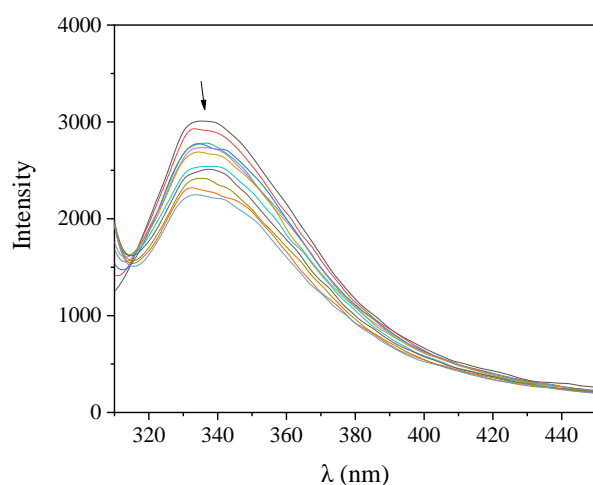


AuNRs@PEG

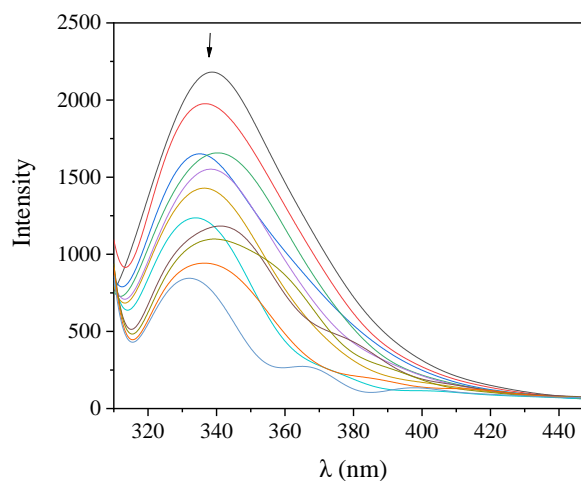


AuNRs@PEG@NAP

Figure S6. Stern-Volmer quenching plot of EB-DNA fluorescence for AuNRs@PEG and AuNRs@PEG@NAP.

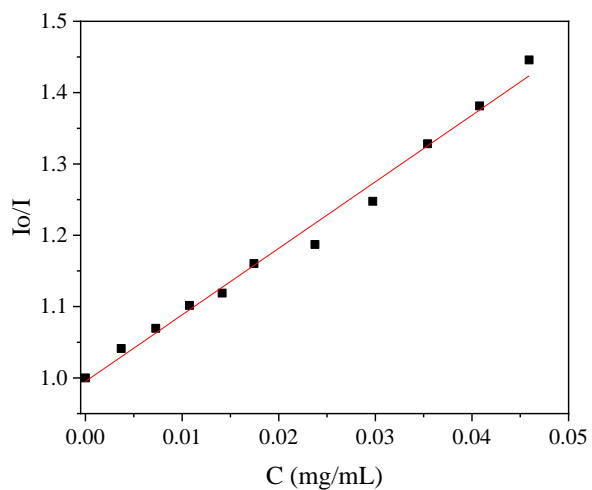


AuNRs@PEG

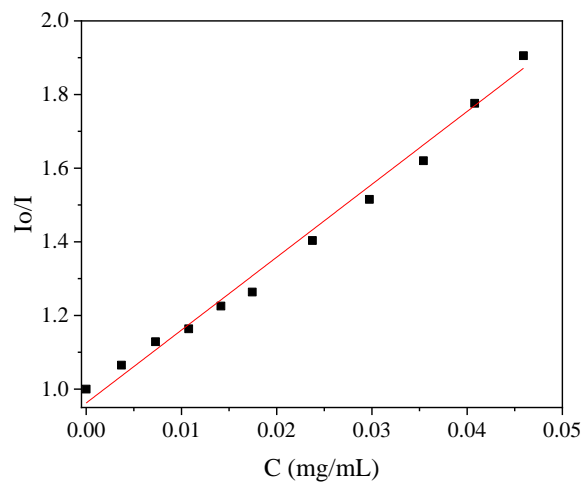


AuNRs@PEG@NAP

Figure S7. Fluorescence emission spectra ($\lambda_{\text{excitation}} = 295 \text{ nm}$) for HSA ($[\text{HSA}] = 3 \mu\text{M}$) in buffer solution (150 mM NaCl and 15 mM trisodium citrate at pH 7.0) in the absence and presence of increasing amounts of AuNRs@PEG and AuNRs@PEG@NAP. The arrow shows the changes of intensity upon increasing amounts of AuNRs@PEG and AuNRs@PEG@NAP.

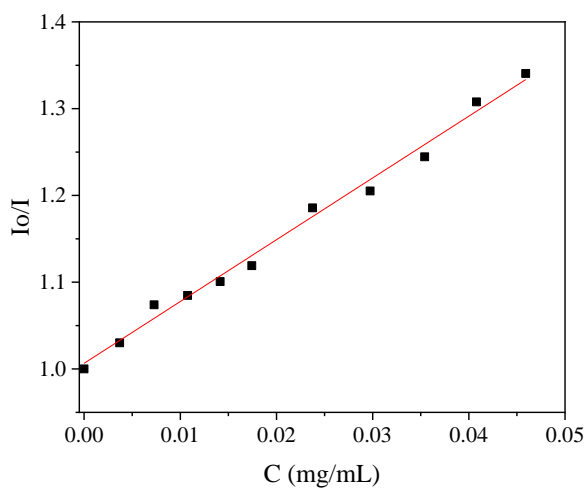


AuNRs@PEG

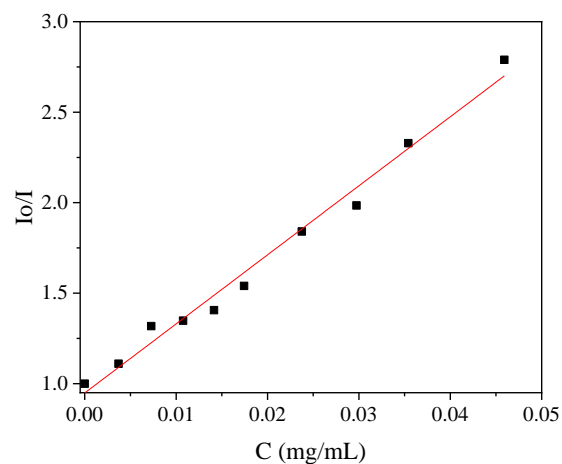


AuNRs@PEG@NAP

Figure S8. Stern-Volmer quenching plot of BSA fluorescence for AuNRs@PEG and AuNRs@PEG@NAP.

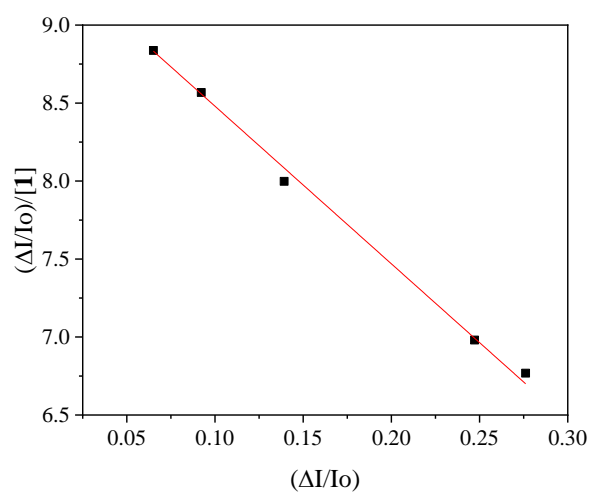


AuNRs@PEG

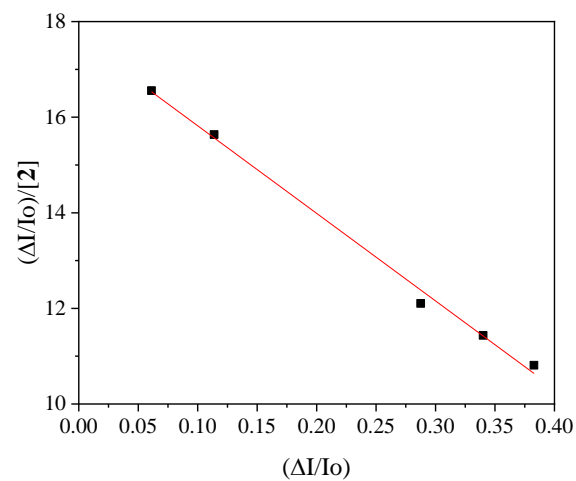


AuNRs@PEG@NAP

Figure S9. Stern-Volmer quenching plot of HSA fluorescence for AuNRs@PEG and AuNRs@PEG@NAP.

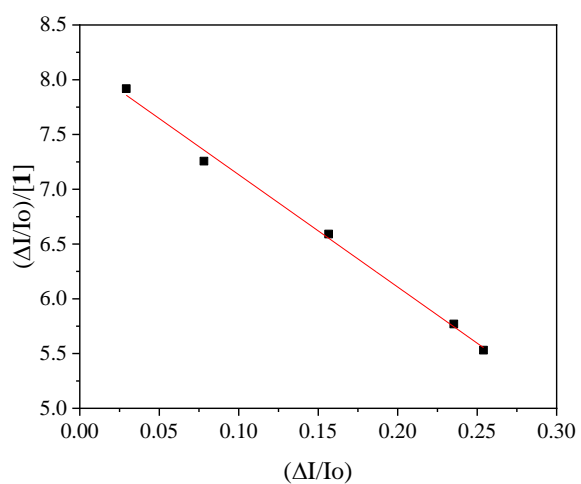


AuNRs@PEG

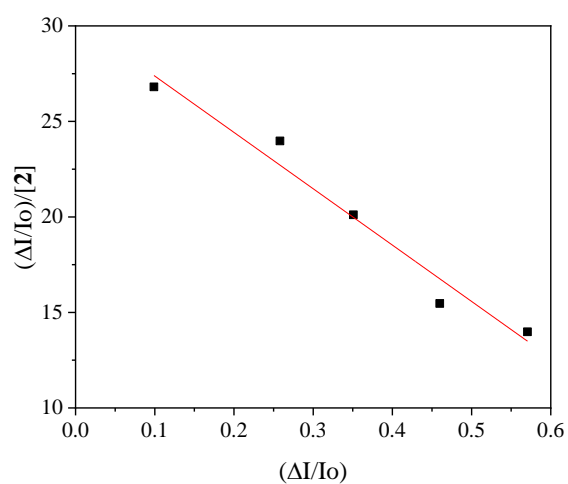


AuNRs@PEG@NAP

Figure S10. Scatchard plot of BSA for AuNRs@PEG (1) and AuNRs@PEG@NAP (2).



AuNRs@PEG



AuNRs@PEG@NAP

Figure S11. Scatchard plot of HSA for AuNRs@PEG (1) and AuNRs@PEG@NAP (2).

S1 Interaction studies with CT DNA

The interaction of the AuNRs with CT DNA was studied by UV-vis spectroscopy at room temperature and variable-temperatures, and *via* competitive studies with EB by fluorescence emission spectroscopy.

S1.1 Binding study with CT DNA by UV-vis spectroscopy

The interaction of the AuNRs with CT DNA has been studied by UV-vis spectroscopy in order to investigate the possible binding modes to CT DNA and to calculate the DNA-binding constants (K_b). The K_b constants (in M^{-1}) were determined by the Wolfe-Shimer equation (eq. S1) [1] and the plots $[DNA]/(\epsilon_A - \epsilon_f)$ *versus* $[DNA]$ using the UV-vis spectra of the AuNRs (0.25 mg/mL) recorded in the presence of increasing concentrations of CT DNA. According to the Wolfe-Shimer equation (equation S1):

$$\frac{[DNA]}{(\epsilon_A - \epsilon_f)} = \frac{[DNA]}{(\epsilon_b - \epsilon_f)} + \frac{1}{K_b(\epsilon_b - \epsilon_f)} \quad (\text{equation S1})$$

where $[DNA]$ is the concentration of DNA in base pairs, $\epsilon_A = A_{\text{obsd}}/[\text{compound}]$, ϵ_f = the extinction coefficient for the free AuNR, and ϵ_b = the extinction coefficient for the AuNR in the fully bound form. K_b is given by the ratio of slope to the y intercept in plots $[DNA]/(\epsilon_A - \epsilon_f)$ *versus* $[DNA]$.

S1.2 Determination of the thermodynamic parameters

In order to determine the thermodynamics parameters enthalpy change (ΔH), entropy change (ΔS) and ΔG , the DNA-binding constants of the AuNRs were determined for three different temperatures (291 K, 300 K and 310 K) as described in section S1.1 with equation S1 and the corresponding plots.

The enthalpy change (ΔH) and the entropy change (ΔS) can be calculated from the van't Hoff equation (equation S2):

$$\ln(K_b) = -\frac{\Delta H}{RT} + \frac{\Delta S}{R} \quad (\text{equation S2})$$

where ΔH and ΔS can be determined from the plot of $\ln(K_b)$ *versus* $(1/T)$, where $-\Delta H/R$ is the slope of the fitting line and $\Delta S/R$ is the intercept (R is the universal gas constant). In addition, ΔG may be obtained from the Gibb's-Helmholtz equation (equation S3):

$$\Delta G = \Delta H - T \cdot \Delta S \quad (\text{equation S3})$$

S1.3 EB-displacement studies

The competition of the AuNRs with EB was investigated by fluorescence emission spectroscopy in order to examine whether the AuNRs may displace EB from its DNA-EB adduct. The CT DNA-EB adduct was formed by pre-treating 20 μM EB and 26 μM CT DNA

in buffer (150 mM NaCl and 15 mM trisodium citrate at pH = 7.0). The possible displacement of EB by the AuNRs and subsequently their intercalating effect was studied by the stepwise addition of a certain amount of the solution of each AuNR into the solution of the CT DNA-EB adduct. The solutions were excited at 540 nm and the emission was monitored from 550-700 nm with $\lambda_{\text{max}} = 592\text{-}595$ nm and the effect of the addition of each AuNR to the CT-DNA EB solution was recorded. The AuNRs did not display any fluorescence emission bands at room temperature in solution or in the presence of CT DNA or EB under the same experimental conditions ($\lambda_{\text{excitation}} = 540$ nm); therefore, the observed quenching of the EB-DNA solution may be attributed to the displacement of EB from its EB-DNA adduct.

The Stern-Volmer constants (K_{SV} , in M^{-1}) were calculated according by the linear Stern-Volmer equation (equation S4) [2] and the respective plots I_0/I versus [AuNR].

$$\frac{I_0}{I} = 1 + k_q \tau_0 [Q] = 1 + K_{\text{SV}} [Q] \quad (\text{equation S4})$$

where I_0 and I are the emission intensities of the EB-DNA solution in the absence and the presence of the AuNRs, respectively, τ_0 = the average lifetime of the emitting system without the quencher and k_q = the quenching constant. Taking $\tau_0 = 23$ ns as the fluorescence lifetime of the EB-DNA adduct [3], the quenching constants (k_q , in $\text{M}^{-1}\text{s}^{-1}$) of the AuNRs were calculated according to equation S5 [2]:

$$K_{\text{SV}} = k_q \tau_0 \quad (\text{equation S5})$$

S2 DNA–cleavage experiments

The reaction mixtures (20 μL) containing supercoiled circular pBR322 DNA stock solution (Form I, 50 μM /base pair, ~ 500 ng), AuNRs, and Tris buffer (25 μM , pH 6.8) in Pyrex vials were incubated for 30 min at 37 $^{\circ}\text{C}$, and was centrifuged under aerobic conditions at room temperature.

After addition of the gel–loading buffer [6X Orange DNA Loading Dye 10 mM Tris–HCl (pH 7.6), 0.15% orange G, 0.03% xylene cyanol FF, 60% glycerol, and 60 mM EDTA, by Fermentas], the reaction mixtures were loaded on a 1% agarose gel with EB staining. The electrophoresis tank was attached to a power supply at a constant current (65 V for 1 h). The gel was visualized by 312 nm UV transilluminator and photographed by an FB–PBC–34 camera vilberlourmat. Quantification of DNA–cleaving activities was performed by integration of the optical density as a function of the band area using the program “Image J” available at the site <http://rsb.info.nih.gov/ij/download.html>.

The ss% and ds% damages were calculated according to the equations S6 and S7:

$$\text{ss\%} = \frac{\text{FormII}}{(\text{FormI} + \text{FormII} + \text{FormIII})} \times 100 \quad (\text{equation S6})$$

$$\text{ds\%} = \frac{\text{FormIII}}{(\text{FormI} + \text{FormII} + \text{FormIII})} \times 100 \quad (\text{equation S7})$$

where, as Form II we consider Form II of each series minus Form II of the irradiated control DNA and as Form I, we consider Form I of each series. The amount of supercoiled DNA was multiplied by factor of 1.43 to account for reduced EB intercalation into supercoiled DNA [4].

S3 Albumin-binding studies

In order to investigate if the AuNRs can bind to carrier proteins like serum albumins, we carried out albumin-binding study by tryptophan fluorescence quenching experiments using bovine serum albumin (BSA, 3 μ M) or human serum albumin (HSA, 3 μ M) in buffer (containing 15 mM trisodium citrate and 150 mM NaCl at pH 7.0). The quenching of the emission intensity of tryptophan residues of BSA at 343 nm was monitored using the AuNRs as quenchers with increasing concentration [2]. The fluorescence emission spectra of the AuNRs were also recorded with $\lambda_{ex} = 295$ nm. No fluorescence emission band was recorded and thus it was not related to the AuNRs. The influence of the inner-filter effect [5] on the measurements was evaluated by equation S8.

$$I_{corr} = I_{meas} \times 10^{\frac{\varepsilon(\lambda_{exc})cd}{2}} \times 10^{\frac{\varepsilon(\lambda_{em})cd}{2}} \quad (\text{equation S8})$$

where I_{corr} = corrected intensity, I_{meas} = the measured intensity, c = the concentration of the quencher, d = the cuvette (1 cm), $\varepsilon(\lambda_{exc})$ and $\varepsilon(\lambda_{em})$ = the ε of the quencher at the excitation and the emission wavelength, respectively, as calculated from the UV-vis spectra of the AuNRs [5].

The Stern-Volmer and Scatchard graphs were used to study the interaction of the compounds with serum albumins. According to Stern-Volmer quenching equation (eq. S2), where I_0 = the initial tryptophan fluorescence intensity of SA, I = the tryptophan fluorescence intensity of SA after the addition of the quencher, k_q = the quenching constant, K_{SV} = the Stern-Volmer constant, τ_0 = the average lifetime of SA without the quencher, and, taking as fluorescence lifetime (τ_0) of tryptophan in SA at around 10^{-8} s [2], K_{SV} can be obtained by the slope of the diagram I_0/I versus $[AuNR]$ (Stern-Volmer plots), and subsequently the quenching (k_q) may be calculated from eq. S3.

From the Scatchard equation (equation S9):

$$\frac{\Delta I/I_0}{[Q]} = nK - K \frac{\Delta I}{I_0} \quad (\text{equation S9})$$

where n is the number of binding sites per albumin and K is the SA-binding constant (K , in M^{-1}) is calculated from the slope in plots $(\Delta I/I_0)/[AuNR]$ versus $\Delta I/I_0$ and n is given by the ratio of y intercept to the slope [2,6].

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